



EXPRESS MAIL NO: EL615485148US

## SEQUENCE LISTING

<110> Itoh, Nobuyuki  
Kavanaugh, W. Michael

<120> HUMAN FGF-23 GENE AND GENE EXPRESSION  
PRODUCTS

<130> PP-17150.001/201130.40901

<140> 09/801,968

<141> 2001-03-07

<160> 46

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 756

<212> DNA

<213> Mus musculus

<400> 1

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cacctgtaca	cggctacagc	caggaccagc	tatcacctac	agatccatag	ggatgggtcat	180
gtagatggca	ccccccatca	gaccatctac	agtgccctga	tgattacatc	agaggacgcc	240
ggctctgtgg	tgataacagg	agccatgact	cgaaggttcc	tttgtatgga	tctccacggc	300
aacatttttg	gatcgcttca	cttcagccca	gagaattgca	agttccgcca	gtggacgctg	360
gagaatggct	atgacgtcta	cttgctgcag	aagcatcact	acctggtgag	cctggggccgc	420
gccaagcgca	ttttccagcc	gggcaccaac	ccgccgccct	tctcccagtt	cctgggtcgc	480
aggaacgagg	tcccgtgct	gcacttctac	actgttcgcc	cacggcgcca	cacgcgcagc	540
gccgaggacc	cacccgagcg	cgacccactg	aacgtgtca	agccgcggcc	ccgcgccacg	600
cctgtgcctg	tatcctgctc	tcgcgagctg	ccgagcgag	aggaagggtg	ccccgcagcc	660
agcgatcctc	tggtgggtgct	gcgcagaggc	cgtggagatg	ctcgcggggg	cgcgggaggc	720
gcggataggt	gtcgcccctt	tcccaggttc	gtctag			756

<210> 2

<211> 251

<212> PRT

<213> Mus musculus

<400> 2

Met	Leu	Gly	Thr	Cys	Leu	Arg	Leu	Leu	Val	Gly	Val	Leu	Cys	Thr	Val
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Cys	Ser	Leu	Gly	Thr	Ala	Arg	Ala	Tyr	Pro	Asp	Thr	Ser	Pro	Leu	Leu
		20						25					30		
Gly	Ser	Asn	Trp	Gly	Ser	Leu	Thr	His	Leu	Tyr	Thr	Ala	Thr	Ala	Arg

Thr Ser Tyr His Leu Gln Ile His Arg Asp Gly His Val Asp Gly Thr  
 50 55 60  
 Pro His Gln Thr Ile Tyr Ser Ala Leu Met Ile Thr Ser Glu Asp Ala  
 65 70 75 80  
 Gly Ser Val Val Ile Thr Gly Ala Met Thr Arg Arg Phe Leu Cys Met  
 85 90 95

Asp Leu His Gly Asn Ile Phe Gly Ser Leu His Phe Ser Pro Glu Asn  
 100 105 110  
 Cys Lys Phe Arg Gln Trp Thr Leu Glu Asn Gly Tyr Asp Val Tyr Leu  
 115 120 125  
 Ser Gln Lys His His Tyr Leu Val Ser Leu Gly Arg Ala Lys Arg Ile  
 130 135 140  
 Phe Gln Pro Gly Thr Asn Pro Pro Phe Ser Gln Phe Leu Ala Arg  
 145 150 155 160  
 Arg Asn Glu Val Pro Leu Leu His Phe Tyr Thr Val Arg Pro Arg Arg  
 165 170 175  
 His Thr Arg Ser Ala Glu Asp Pro Pro Glu Arg Asp Pro Leu Asn Val  
 180 185 190  
 Leu Lys Pro Arg Pro Arg Ala Thr Pro Val Pro Val Ser Cys Ser Arg  
 195 200 205  
 Glu Leu Pro Ser Ala Glu Glu Gly Gly Pro Ala Ala Ser Asp Pro Leu  
 210 215 220  
 Gly Val Leu Arg Arg Gly Arg Gly Asp Ala Arg Gly Gly Ala Gly Gly  
 225 230 235 240  
 Ala Asp Arg Cys Arg Pro Phe Pro Arg Phe Val  
 245 250

<210> 3  
 <211> 756  
 <212> DNA  
 <213> Homo sapiens

<400> 3  
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 cacctgtaca cagccacagc caggaacagc taccacctgc agatccacaa gaatggccat 180  
 gtggatggcg caccatca gaccatctac agtgccctga tgatcagatc agaggatgct 240  
 ggctttgtgg tgattacagg tgtgatgagc agaagatacc tctgcatgga tttcagaggc 300  
 aacatttttg gatcacacta tttcgacccg gagaactgca ggttccaaca ccagacgctg 360  
 gaaaacgggt acgacgtcta ccactctcct cagtatcact tcttggtcag tctgggccgg 420  
 gcgaagagag ccttctctgcc aggcattgaac ccacccccgt actccagtt cctgtcccgg 480  
 aggaacgaga tccccctaatt tcaactcaac acccccatc caccggcgga caccgggagc 540  
 gccgaggacg actcggagcg ggacccccctg aacgtgctga agccccgggc ccggatgacc 600  
 ccggccccgg cctcctgttc acaggagctc ccgagcgccg aggacaacag cccgatggcc 660  
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<210> 4  
 <211> 251  
 <212> PRT  
 <213> Homo sapiens

<400> 4  
 Met Leu Gly Ala Arg Leu Arg Leu Trp Val Cys Ala Leu Cys Ser Val

1				5					10					15			
Cys	Ser	Met	Ser	Val	Leu	Arg	Ala	Tyr	Pro	Asn	Ala	Ser	Pro	Leu	Leu		
			20					25					30				
Gly	Ser	Ser	Trp	Gly	Gly	Leu	Ile	His	Leu	Tyr	Thr	Ala	Thr	Ala	Arg		
		35					40					45					
Asn	Ser	Tyr	His	Leu	Gln	Ile	His	Lys	Asn	Gly	His	Val	Asp	Gly	Ala		
	50					55				60							
Pro	His	Gln	Thr	Ile	Tyr	Ser	Ala	Leu	Met	Ile	Arg	Ser	Glu	Asp	Ala		
65				70					75					80			
Gly	Phe	Val	Val	Ile	Thr	Gly	Val	Met	Ser	Arg	Arg	Tyr	Leu	Cys	Met		
			85					90					95				
Asp	Phe	Arg	Gly	Asn	Ile	Phe	Gly	Ser	His	Tyr	Phe	Asp	Pro	Glu	Asn		
		100					105					110					
Cys	Arg	Phe	Gln	His	Gln	Thr	Leu	Glu	Asn	Gly	Tyr	Asp	Val	Tyr	His		
	115					120						125					
Ser	Pro	Gln	Tyr	His	Phe	Leu	Val	Ser	Leu	Gly	Arg	Ala	Lys	Arg	Ala		
	130				135					140							
Phe	Leu	Pro	Gly	Met	Asn	Pro	Pro	Pro	Tyr	Ser	Gln	Phe	Leu	Ser	Arg		
145				150					155					160			
Arg	Asn	Glu	Ile	Pro	Leu	Ile	His	Phe	Asn	Thr	Pro	Ile	Pro	Arg	Arg		
			165				170							175			
His	Thr	Arg	Ser	Ala	Glu	Asp	Asp	Ser	Glu	Arg	Asp	Pro	Leu	Asn	Val		
		180				185						190					
Leu	Lys	Pro	Arg	Ala	Arg	Met	Thr	Pro	Ala	Pro	Ala	Ser	Cys	Ser	Gln		
	195					200						205					
Glu	Leu	Pro	Ser	Ala	Glu	Asp	Asn	Ser	Pro	Met	Ala	Ser	Asp	Pro	Leu		
	210				215						220						
Gly	Val	Val	Arg	Gly	Gly	Arg	Val	Asn	Thr	His	Ala	Gly	Gly	Thr	Gly		
225				230					235					240			
Pro	Glu	Gly	Cys	Arg	Pro	Phe	Ala	Lys	Phe	Ile							
			245			250											

<210> 5  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Sense PCR primer

<400> 5  
 agcaccagcc actcagagca

20

<210> 6  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Antisense PCR primer

<400> 6  
 cttccagcga ccctagatga

20

<210> 7  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence

<220>

<223> Sense primer for mouse FGF-23

<400> 7  
 ctgatgatta catcagagga c 21

<210> 8  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence

<220>

<223> Antisense primer for mouse FGF-23

<400> 8  
 caccaggtag tgatgcttct 20

<210> 9  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence

<220>

<223> Antisense primer for mouse FGF-23

<400> 9  
 atccatacaa aggaaccttc g 21

<210> 10  
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 <212> DNA  
 <213> Artificial Sequence

<220>

<223> adaptor primer

<400> 10  
 ccatcctaata acgactcact atagggc 27

<210> 11  
 <211> 23  
 <212> DNA  
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<220>

<223> adaptor primer

<400> 11  
 actcactata gggctcgagc ggc 23

<210> 12  
 <211> 20  
 <212> DNA  
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<220>  
 <223> Sense primer for mouse FGF-23.

<400> 12  
 actcagtgtgt gtgcaatgtct 20

<210> 13  
 <211> 20  
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 <213> Artificial Sequence

<220>  
 <223> Antisense primer for mouse FGF-23

<400> 13  
 gacctagacg aacctgggaa 20

<210> 14  
 <211> 216  
 <212> PRT  
 <213> Homo sapiens

<400> 14  
 Met Arg Ser Gly Cys Val Val Val His Val Trp Ile Leu Ala Gly Leu  
 1 5 10 15  
 Trp Leu Ala Val Ala Gly Arg Pro Leu Ala Phe Ser Asp Ala Gly Pro  
 20 25 30  
 His Val His Tyr Gly Trp Gly Asp Pro Ile Arg Leu Arg His Leu Tyr  
 35 40 45  
 Thr Ser Gly Pro His Gly Leu Ser Ser Cys Phe Leu Arg Ile Arg Ala  
 50 55 60  
 Asp Gly Val Val Asp Cys Ala Arg Gly Gln Ser Ala His Ser Leu Leu  
 65 70 75 80  
 Glu Ile Lys Ala Val Ala Leu Arg Thr Val Ala Ile Lys Gly Val His  
 85 90 95  
 Ser Val Arg Tyr Leu Cys Met Gly Ala Asp Gly Lys Met Gln Gly Leu  
 100 105 110  
 Leu Gln Tyr Ser Glu Glu Asp Cys Ala Phe Glu Glu Glu Ile Arg Pro  
 115 120 125  
 Asp Gly Tyr Asn Val Tyr Arg Ser Glu Lys His Arg Leu Pro Val Ser  
 130 135 140  
 Leu Ser Ser Ala Lys Gln Arg Gln Leu Tyr Lys Asn Arg Gly Phe Leu  
 145 150 155 160  
 Pro Leu Ser His Phe Leu Pro Met Leu Pro Met Val Pro Glu Glu Pro  
 165 170 175  
 Glu Asp Leu Arg Gly His Leu Glu Ser Asp Met Phe Ser Ser Pro Leu  
 180 185 190  
 Glu Thr Asp Ser Met Asp Pro Phe Gly Leu Val Thr Gly Leu Glu Ala

195 200 205  
 Val Arg Ser Pro Ser Phe Glu Lys  
 210 215

<210> 15  
 <211> 209

<212> PRT  
 <213> Homo sapiens

<400> 15

Met Asp Ser Asp Glu Thr Gly Phe Glu His Ser Gly Leu Trp Val Ser  
 1 5 10 15  
 Val Leu Ala Gly Leu Leu Leu Gly Ala Cys Gln Ala His Pro Ile Pro  
 20 25 30  
 Asp Ser Ser Pro Leu Leu Gln Phe Gly Gly Gln Val Arg Gln Arg Tyr  
 35 40 45  
 Leu Tyr Thr Asp Asp Ala Gln Gln Thr Glu Ala His Leu Glu Ile Arg  
 50 55 60  
 Glu Asp Gly Thr Val Gly Gly Ala Ala Asp Gln Ser Pro Glu Ser Leu  
 65 70 75 80  
 Leu Gln Leu Lys Ala Leu Lys Pro Gly Val Ile Gln Ile Leu Gly Val  
 85 90 95  
 Lys Thr Ser Arg Phe Leu Cys Gln Arg Pro Asp Gly Ala Leu Tyr Gly  
 100 105 110  
 Ser Leu His Phe Asp Pro Glu Ala Cys Ser Phe Arg Glu Leu Leu Leu  
 115 120 125  
 Glu Asp Gly Tyr Asn Val Tyr Gln Ser Glu Ala His Gly Leu Pro Leu  
 130 135 140  
 His Leu Pro Gly Asn Lys Ser Pro His Arg Asp Pro Ala Pro Arg Gly  
 145 150 155 160  
 Pro Ala Arg Phe Leu Pro Leu Pro Gly Leu Pro Pro Ala Leu Pro Glu  
 165 170 175  
 Pro Pro Gly Ile Leu Ala Pro Gln Pro Pro Asp Val Gly Ser Ser Asp  
 180 185 190  
 Pro Leu Ser Met Val Gly Pro Ser Gln Gly Arg Ser Pro Ser Tyr Ala  
 195 200 205  
 Ser

<210> 16  
 <211> 10  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Residues which can be incorporated to allow myc  
 monoclonal antibody-based affinity purification.

<400> 16

Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu  
 1 5 10

<210> 17  
 <211> 5

<212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Preferred thrombin cleavage site.

<400> 17  
 Leu Val Pro Arg Gly  
 1 5

<210> 18  
 <211> 10  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Residues that bind to paramagnetic streptavidin  
 beads which facilitates purification of molecules.

<400> 18  
 Ser Ala Trp Arg His Pro Gln Phe Gly Gly  
 1 5 10

<210> 19  
 <211> 14  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Oligopeptide used for the production of an  
 antibody to FGF-23 protein. (residues 175-189 of  
 SEQ ID NO:4)

<400> 19  
 Arg Arg His Thr Arg Ser Ala Glu Asp Asp Ser Glu Arg Asp  
 1 5 10

<210> 20  
 <211> 17  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Oligopeptide used for the production of an  
 antibody to FGF-23 protein. (residues 51-67 of  
 SEQ ID NO:4)

<400> 20  
 Tyr His Leu Gln Ile His Lys Asn Gly His Val Asp Gly Ala Pro His  
 1 5 10 15  
 Gln

<210> 21





Ser Ser Asp  
35

<210> 25  
<211> 34  
<212> PRT

<213> Homo sapiens

<400> 25  
Gln Met Tyr Val Ala Leu Asn Gly Lys Gly Ala Pro Arg Arg Gly Gln  
1 5 10 15  
Lys Thr Arg Arg Lys Asn Thr Ser Ala His Phe Leu Pro Met Val Val  
20 25 30  
His Ser

<210> 26  
<211> 56  
<212> PRT  
<213> Homo sapiens

<400> 26  
Ala Trp Tyr Leu Gly Leu Asp Lys Glu Gly Gln Val Met Lys Gly Asn  
1 5 10 15  
Arg Val Lys Lys Thr Lys Ala Ala Ala His Phe Leu Pro Lys Leu Leu  
20 25 30  
Glu Val Ala Met Tyr Gln Glu Pro Ser Leu His Ser Val Pro Glu Ala  
35 40 45  
Ser Pro Ser Ser Pro Pro Ala Pro  
50 55

<210> 27  
<211> 72  
<212> PRT  
<213> Homo sapiens

<400> 27  
Ala Trp Phe Leu Gly Leu Asn Lys Glu Gly Gln Ile Met Lys Gly Asn  
1 5 10 15  
Arg Val Lys Lys Thr Lys Pro Ser Ser His Phe Val Pro Lys Pro Ile  
20 25 30  
Glu Val Cys Met Tyr Arg Glu Pro Ser Leu His Glu Ile Gly Glu Lys  
35 40 45  
Gln Gly Arg Ser Arg Lys Ser Ser Gly Thr Pro Thr Met Asn Gly Gly  
50 55 60  
Lys Val Val Asn Gln Asp Ser Thr  
65 70

<210> 28  
<211> 78  
<212> PRT  
<213> Homo sapiens

<400> 28

Gly	Trp	Tyr	Leu	Gly	Leu	Asn	Lys	Glu	Gly	Glu	Ile	Met	Lys	Gly	Asn
1				5					10					15	
His	Val	Lys	Lys	Asn	Lys	Pro	Ala	Ala	His	Phe	Leu	Pro	Lys	Pro	Leu
			20					25					30		
Lys	Val	Ala	Met	Tyr	Lys	Glu	Pro	Ser	Leu	His	Asp	Leu	Thr	Glu	Phe
		35					40					45			

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Ser	Arg	Ser	Gly	Ser	Gly	Thr	Pro	Thr	Lys	Ser	Arg	Ser	Val	Ser	Gly
	50					55					60				
Val	Leu	Asn	Gly	Gly	Lys	Ser	Met	Ser	His	Asn	Glu	Ser	Thr		
65					70					75					

<210> 29  
 <211> 78  
 <212> PRT  
 <213> Homo sapiens

Ala	Trp	Phe	Leu	Gly	Leu	Asn	Lys	Glu	Gly	Gln	Ala	Met	Lys	Gly	Asn
1				5					10					15	
Arg	Val	Lys	Lys	Thr	Lys	Pro	Ala	Ala	His	Phe	Leu	Pro	Lys	Pro	Leu
			20					25					30		
Glu	Val	Ala	Met	Tyr	Arg	Glu	Pro	Ser	Leu	His	Asp	Val	Gly	Glu	Thr
		35					40					45			
Val	Pro	Lys	Pro	Gly	Val	Thr	Pro	Ser	Lys	Ser	Thr	Ser	Ala	Ser	Ala
	50					55					60				
Ile	Met	Asn	Gly	Gly	Lys	Pro	Val	Asn	Lys	Ser	Lys	Thr	Thr		
65					70					75					

<210> 30  
 <211> 78  
 <212> PRT  
 <213> Homo sapiens

Ala	Trp	Phe	Leu	Gly	Leu	Asn	Lys	Glu	Gly	Gln	Ala	Met	Lys	Gly	Asn
1				5					10					15	
Arg	Val	Lys	Lys	Thr	Lys	Pro	Ala	Ala	His	Phe	Leu	Pro	Lys	Pro	Leu
			20					25					30		
Glu	Val	Ala	Met	Tyr	Arg	Glu	Pro	Ser	Leu	His	Asp	Val	Gly	Glu	Thr
		35					40					45			
Val	Pro	Lys	Pro	Gly	Val	Thr	Pro	Ser	Lys	Ser	Thr	Ser	Ala	Ser	Ala
	50					55					60				
Ile	Met	Asn	Gly	Gly	Lys	Pro	Val	Asn	Lys	Ser	Lys	Thr	Thr		
65					70					75					

<210> 31  
 <211> 48  
 <212> PRT  
 <213> Homo sapiens

Gln	Tyr	Tyr	Val	Ala	Leu	Asn	Lys	Asp	Gly	Ser	Pro	Arg	Glu	Gly	Tyr
1				5					10					15	
Arg	Thr	Lys	Arg	His	Gln	Lys	Phe	Thr	His	Phe	Leu	Pro	Arg	Pro	Val

Asp Pro Ser Lys Leu Pro Ser Met Ser Arg Asp Leu Phe His Tyr Arg  
 35 40 45

<210> 32

<211> 68

<212> PRT

<213> Homo sapiens

<400> 32

Trp Phe Met Ala Phe Thr Arg Gln Gly Arg Pro Arg Gln Ala Ser Arg  
 1 5 10 15  
 Ser Arg Gln Asn Gln Arg Glu Ala His Phe Ile Lys Arg Leu Tyr Gln  
 20 25 30  
 Gly Gln Leu Pro Phe Pro Asn His Ala Glu Lys Gln Lys Gln Phe Glu  
 35 40 45  
 Phe Val Gly Ser Ala Pro Thr Arg Arg Thr Lys Arg Thr Arg Arg Pro  
 50 55 60  
 Gln Pro Leu Thr  
 65

<210> 33

<211> 59

<212> PRT

<213> Homo sapiens

<400> 33

Trp Tyr Val Gly Phe Thr Lys Lys Gly Arg Pro Arg Lys Gly Pro Lys  
 1 5 10 15  
 Thr Arg Glu Asn Gln Gln Asp Val His Phe Met Lys Arg Tyr Pro Lys  
 20 25 30  
 Gly Gln Pro Glu Leu Gln Lys Pro Phe Lys Tyr Thr Thr Val Thr Lys  
 35 40 45  
 Arg Ser Arg Arg Ile Arg Pro Thr His Pro Ala  
 50 55

<210> 34

<211> 76

<212> PRT

<213> Homo sapiens

<400> 34

Leu Pro Val Ser Leu Ser Ser Ala Lys Gln Arg Gln Leu Tyr Lys Asn  
 1 5 10 15  
 Arg Gly Phe Leu Pro Leu Ser His Phe Leu Pro Met Leu Pro Met Val  
 20 25 30  
 Pro Glu Glu Pro Glu Asp Leu Arg Gly His Leu Glu Ser Asp Met Phe  
 35 40 45  
 Ser Ser Pro Leu Glu Thr Asp Ser Met Asp Pro Phe Gly Leu Val Thr  
 50 55 60  
 Gly Leu Glu Ala Val Arg Ser Pro Ser Phe Glu Lys  
 65 70 75

<210> 35

<211> 33  
 <212> PRT  
 <213> Homo sapiens

<400> 35  
 Trp Tyr Val Ala Leu Lys Arg Thr Gly Gln Tyr Lys Leu Gly Ser Lys  
 1 5 10 15  
 Thr Gly Pro Gly Gln Lys Ala Ile Leu Phe Leu Pro Met Ser Ala Lys  
 20 25 30  
 Ser

<210> 36  
 <211> 68  
 <212> PRT  
 <213> Homo sapiens

<400> 36  
 Leu Pro Leu His Leu Pro Gly Asn Lys Ser Pro His Arg Asp Pro Ala  
 1 5 10 15  
 Pro Arg Gly Pro Ala Arg Phe Leu Pro Leu Pro Gly Leu Pro Pro Ala  
 20 25 30  
 Leu Pro Glu Pro Pro Gly Ile Leu Ala Pro Gln Pro Pro Asp Val Gly  
 35 40 45  
 Ser Ser Asp Pro Leu Ser Met Val Gly Pro Ser Gln Gly Arg Ser Pro  
 50 55 60  
 Ser Tyr Ala Ser  
 65

<210> 37  
 <211> 88  
 <212> PRT  
 <213> Homo sapiens

<400> 37  
 Leu Trp Tyr Val Ser Val Asn Gly Lys Gly Arg Pro Arg Arg Gly Phe  
 1 5 10 15  
 Lys Thr Arg Arg Thr Gln Lys Ser Ser Leu Phe Leu Pro Arg Val Leu  
 20 25 30  
 Asp His Arg Asp His Glu Met Val Arg Gln Leu Gln Ser Gly Leu Pro  
 35 40 45  
 Arg Pro Pro Gly Lys Gly Val Gln Pro Arg Arg Arg Gln Lys Gln  
 50 55 60  
 Ser Pro Asp Asn Leu Glu Pro Ser His Val Gln Ala Ser Arg Leu Gly  
 65 70 75 80  
 Ser Gln Leu Glu Ala Ser Ala His  
 85

<210> 38  
 <211> 30  
 <212> PRT  
 <213> Homo sapiens

<400> 38

Protein Data Bank

Met Phe Ile Ala Leu Ser Lys Asn Gly Lys Thr Lys Lys Gly Asn Arg  
 1 5 10 15  
 Val Ser Pro Thr Met Lys Val Thr His Phe Leu Pro Arg Leu  
 20 25 30

<210> 39  
 <211> 82  
 <212> PRT  
 <213> Homo sapiens

<400> 39  
 Glu Trp Tyr Val Ala Leu Asn Lys Arg Gly Lys Ala Lys Arg Gly Cys  
 1 5 10 15  
 Ser Pro Arg Val Lys Pro Gln His Ile Ser Thr His Phe Leu Pro Arg  
 20 25 30  
 Phe Lys Gln Ser Glu Gln Pro Glu Leu Ser Phe Thr Val Thr Val Pro  
 35 40 45  
 Glu Lys Lys Lys Pro Pro Ser Pro Ile Lys Pro Lys Ile Pro Leu Ser  
 50 55 60  
 Ala Pro Arg Lys Asn Thr Asn Ser Val Lys Tyr Arg Leu Lys Phe Arg  
 65 70 75 80  
 Phe Gly

<210> 40  
 <211> 30  
 <212> PRT  
 <213> Homo sapiens

<400> 40  
 Thr Tyr Ile Ala Leu Ser Lys Tyr Gly Arg Val Lys Arg Gly Ser Lys  
 1 5 10 15  
 Val Ser Pro Ile Met Thr Val Thr His Phe Leu Pro Arg Ile  
 20 25 30

<210> 41  
 <211> 33  
 <212> PRT  
 <213> Homo sapiens

<400> 41  
 Glu Met Phe Val Ala Leu Asn Gln Lys Gly Ile Pro Val Arg Gly Lys  
 1 5 10 15  
 Lys Thr Lys Lys Glu Gln Lys Thr Ala His Phe Leu Pro Met Ala Ile  
 20 25 30  
 Thr

<210> 42  
 <211> 67  
 <212> PRT  
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<400> 42

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<223> consensus sequence

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